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OM nucleic - nucleic search, using sw model
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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/Backfiles1.seq:*
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US-08-637-509-3
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US-08-647-33-659A-9
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US-08-634-642-1
US-08-634-642-1
US-08-684-642-1
US-09-058-489-13
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ALIGNMENTS

US-09-008-271A-16 Sequence 16, Application US/09008271A Patent No. 6203979 GENERAL INFORMATION: APPLICANT: Bandman, Olga APPLICANT: Teopifor I. ; SEQUENCE DESCRIPTION: SEQ ID NO: 16 US-09-008-271A-16 INFORMATION FOR SEQ ID NO: 16: CORRESPONDENCE ADDRESS: ADDRESSEE: Incyte Pharmaceuticals, Inc STREET: 3174 Porter Dr NAME: Mohan-Peterson, Sheela REGISTRATION NUMBER: 41.201 REFERENCE/DOCKET NUMBER: PF-0458 US TELECOMMUNICATION INFORMATION: TELEPHONE: 650-855-0555 TELEPHONE: 650-845-4186 OPERATING SYSTEM: DOS SOFTWARE: FASTSEO for Windows Version 2.0 CURRENT APPLICATION DATA: APPLICATION UNMBER: US/09/008,271A FILING DATE: 16-Jan-1998 PRIOR APPLICATION DATA: COMPUTER READABLE FORM: MEDIUM TYPE: Diskette TITLE OF INVENTION: HUMAN PROTEASE MOLECULES NUMBER OF SEQUENCES: 24 APPLICATION NUMBER: <Unknown> FILING DATE: <Unknown> ATTORNEY_AGENT INFORMATION: SEQUENCE CHARACTERISTICS: LENGTH: 2061 base pairs IMMEDIATE SOURCE: COUNTRY: USA ZIP: 94304 COMPUTER: STATE: CA CITY: Palo Alto LIBRARY: LUNGAST01 CLONE: 877617 TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: Guegler, Karl J Corley, Neil C. Tang, Tom Y. Shah, Purvi Hillman, IBM Compatible Henry linear Jenniter L Karl J.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  106 CTCTGCCACCATGACCAAGAAGGTTCAAAAACTTCAGCGAGCAGCTCTGAAGAATCCTGT 165
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                                                                                                                                                                                                                                                                526 CAAGGATTACATCCATCGAGTAGGTCGAACAGCTAGAGCTGGGCGCTCCGGAAAGGCTAT 585
                                                                                          APPLICANT:
APPLICANT:
                                                                                                                                                                                                                           586 TACTTTTGTCACACAGTATGATGTGGAACTCTTTCCAGCGCATAGAACACTTAATTGGGAA
                                                    TITLE OF INVENTION: Lactobacilli Harboring Aggregation and Muc. TITLE OF INVENTION: Binding Genes As Vaccine Delivery Vehicles NUMBER OF SEQUENCES: 2
                                      CORRESPONDENCE ANDRESS:
ADDRESSEE: Standley
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                                     ADDRESSEE:
COUNTRY:
                  Dublin
          Ohio
                            495 Metro Place South,
                                                                                                      Jonsson, Hans
                                                                                                                 Casa
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                                                                                             Mvllstam, Bo
                                        ndley & Gilcrest
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                                                                            Lactobacilli Harboring Aggregation and Mucin
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                                  Suite 210
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CELL TYPE: Unicellular organism US-09-039-773A-1
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INFORMATION FOR SEQ II NO: 1:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
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ORIGINAL SOURCE:
ORGANISM: Lact
STRAIN: 1063
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                                                                                                                                                                                                                                                                                                                                                                       162 aaacccaccaagatccagattgaagctattcctttggccttacaaggtcgtgatatcatt 221
                                                                                                                                                                                                                                                                                                  367 GTTGATACTGAAAATCCCAATATTCAAGCAATTATCATTTCACCAACACGTGAATTAGCG 426
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 4
                                                                                                  487 GTCTATGGTGGGGCAGATATTCGGCGCCCAAATTAAGAGCTTGAAACAACACCCCCAAATT 546
                                                                                                                                                             427 ATCCAGACCCAAGAAGAACFTTATCGTCTAGGTAAAGATAAACATGTTCGCGTGCAGGTA 486
                                                                                                                                                                                        342 tttcagatctcagagcag#ttgaagccctggggtcctctattggagtgcagagtgctgtg 401
                                                                                                                              402 attgtaggtggaattgatt\varphiaatgtctcaatctttggcccttgcaaaaaaaaccacatata 461
                                                                 462 ataatagcaactcctggtcgactgattgaccacttggaaaaatacgaaaggtttcaacttg 521
                                        547 CTCGTGGGGACCCCTGGACG
        522 agagototoaaataottggtфatggatgaagoogacogaataotgaatatggattitgag 581
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792-5536
388 applicable
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches 557; Indels
                                                 GTTACGTGACCATATTAACCGTCATACAG---TTAAACTT 603
      3; Gaps
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